SEQUENCE LISTING

- (1) GENERAL INFORMATION
 - (i) APPLICANT: Elizabeth Frayne
 - (ii) APPLICANT ADRESS: 2027 Galvin Ln. #1
 - (iii) TITLE OF INVENTION: Microbial Production of Nuclease Resistant DNA, RNA, and Oligo Mixtures
 - (iv) NUMBER OF SEQUENCES: 2
 - (v) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Frayne Consultants
 - (B) STREET: 1249 S. Diamond Bar Blvd. #125
 - (C) CITY: Diamond Bar
 - (D) STATE: California
 - (E) COUNTRY: USA
 - (F) ZIP: 91765
 - (iv) COMPUTER READABLE FORM
 - (A) MEDIUM TYPE: Diskette, 3.50 inch 1.2 KB
 - (B) COMPUTER: IBM
 - (C) OPERATING SYSTEM: Windows 98
 - (D) SOFTWARE: Microsoft Word
 - (vii) CURRENT APPLICATION DATA
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION: 435/41,435/71, 435/91, 435/131, 435/154, 435/172, 435/183; 935, 536/27
 - (viii) PRIOR APLICATION DATA:
 - (A) APPLICATION (PROVISIONAL) NUMBER: 60/100,231
 - (B) FILING DATE: 9/14/98
 - (C) APPLICATION (PROVISIONAL) NUMBER: 60/152,371
 - (D) FILING DATE: 9/07/99
 - (viiii) TELECOMUNICATION INFORMATION
 - (A) TELEPHONE: (909)861-2111/860-7415
 - (B) EMAIL: egfrayne@cs.com
- (2) INFORMATION FOR SEQ ID NO.1:
 - (i)SEQUENCE CHRACTERISTICS
 - (A) LENGTH: 22 bp
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE:
 - (A) DESCRIPTION: oligonucleotide
- (iii) HYPOTHETICAL: no
- (iv) ANTI-SENSE: no
- (v) ORIGINAL SOURCE:
 - (A) ORGANSIM: Saccharomyces cerevisiae
 - (B) STRAIN: YFLO39C genomic clone
- (vi) PUBLICATION INFORMATION:
 - (A) AUTHORS: Murakami Y., Naitou M., Hagiwara H., Shibata T., Ozawa M., Sasanuma S.I., Sasanuma M., Tsuchiya Y., Soeda E., Yokoyama K., Yamazaki M., Tashiro H., Eki T.
 - (B) TITLE: Analysis of the nucleotide sequence of chromosome VI from Saccharomyces cerevisiae.
 - (C) JOURNAL: Nature Genet.
 - (D) VOLUME: 10
 - (C) PAGES: 261-268
 - (D) DATE:1995
 - (E) RELEVANT RESIDUES IN SEQ ID NO. 1: 10-31
- (vii) SEQUENCE DESCRIPTION: SEQ ID NO. 1:

GAGGTTGCTG CTTTGGTTAT TG

- (3) INFORMATION FOR SEQ ID NO. 2:
 - (i)SEQUENCE CHRACTERISTICS
 - (E) LENGTH: 22 bp
 - (F) TYPE: nucleic acid
 - (G) STRANDEDNESS: single
 - (H) TOPOLOGY: linear
 - (ii) MOLECULE TYPE:
 - (B) DESCRIPTION: oligonucleotide
 - (iii) HYPOTHETICAL: no
 - (iv) ANTI-SENSE: no
 - v) ORIGINAL SOURCE:
 - (F) ORGANSIM: Saccharomyces cerevisiae
 - (G) STRAIN: YFLO39C genomic clone
 - (vi) PUBLICATION INFORMATION:
 - (A) AUTHORS: Murakami Y., Naitou M., Hagiwara H., Shibata T., Ozawa M., Sasanuma S.I., Sasanuma M., Tsuchiya Y., Soeda E., Yokoyama K., Yamazaki M., Tashiro H., Eki T.